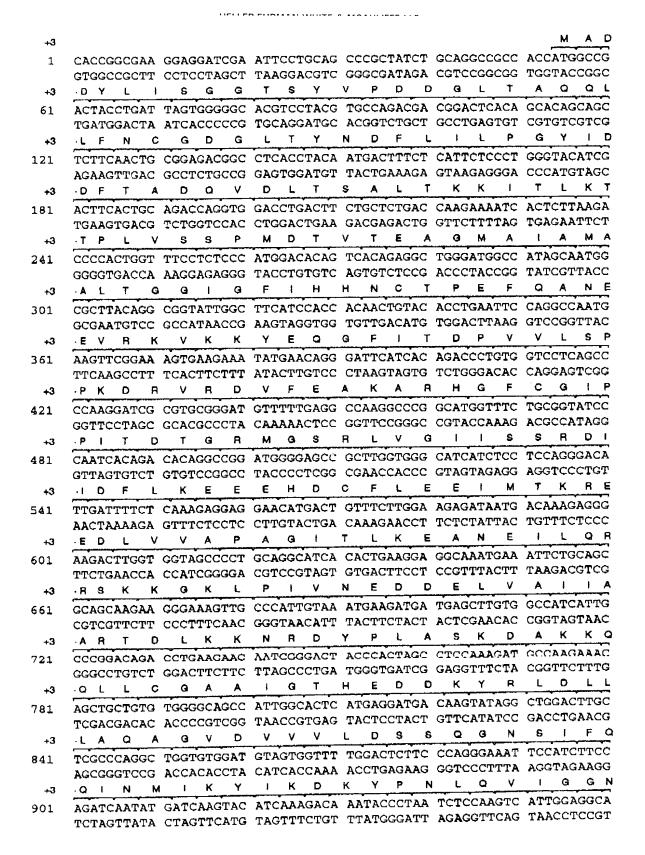
Sheet 1 of 7

ELECTION SYSTEMS FOR GENETICALLY

MODIFIED CELLS

DOCKET NO. 24751-2502

Applicant: Jensen Filed: April 30, 2001





HELLER EHRMAN WHITE & MCAULIFFE LLP Sheet 2 of 7 SELECTION SYSTEMS FOR GENETICALLY

MODIFIED CELLS

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961														AGGT						
	TACA	CCAG	TG	ACG	ACGGG	GTC	CGG	TTC	TTGG	AG	TAA	CTAC	CG	TCCA	CAC	CTA	CGG	GAC	GCC	Ç
+3	∙V G	М	Ģ	S	G	8		С	1	1	Q	E		L	A	С	G	R	P	a C
1021	TGGG	CATG	GG	AAG	rggc'	rcc	ATC	TGC	ATTA	TC	CAG	GAAC	T	GCTG	GCC	TGT	GGG	CGG	CCC	C
	ACCC	GTAC	CC	TTC	ACCG	AGG	TAG	ACG	TAAT	AG	GTC	CTTC	A	CGAC	CGG	ACA	CCC	GCC	GGG	G
+3	·Q A		A		Y	Κ		Y	E	Y		R	R		G	٧		٧	1	
1081	AAGC	AAÇA	GC	AGTO	TAC	AAG	GTG	TAT	GAGT	ΑТ	GCA	ccac	2G	CTTT	GGT	GTT	CCG	GTC.	ATT	G
	TTCG	TTGT	CG	TCAC	ATG	rtc	CAC	ATA	CTCA	TA	CGT	GCC	3C	GAAA	.CCA	CAA	GG¢	ÇAG	TAA	С
+3	· A D	G	Q		Q	N	٧	G	Н	1	Α	ĸ	A	L	A	L	G	Α_	S	Ţ
1141														CTTG						
	GACT	ACCT	CC	TTAC	GTT.	ГТА	CAC	CCA	GTAT	AA	CGC	TTTC	CG	GAAC	CGG	GAA	CC¢	CGG.	AGG	T
+3	·T V	М	М	G	S	L	L	A	Α	T	T	E	Α	P	G	E	Y	F_	F.	S
1201														CCCT						
	GTCA	GTAC	TΑ	CCCG	AGA(GAG	GAC	CGA	CGGT	GG	TGA	CTCC	CG	GGGA	CCA	CTT	ATG	AAG.		
+3	·s D	(3	- 1	R	L	K	К	Υ	R	G	M	G	S	L	D	A	М	D	<u> </u>	H
1261														TCTC						
	GGCT	ACCC	TA	GGCC	GAT	rtc	TTT.	АТА	GCGC	CA	TAC	CCAA	\G	AGAG	CTA	CGG	TAC	CTG'		
+3	·H L	9	S	a	N	R	Y	F	3	E	A	D	K	<u>, l</u>	K	<u> </u>	, A	Q,	G.	
1321														AATC						
	TGGA	GTCG	TC	GGTC	TTG	CT	ATA	AAG	TCAC	TT	CGA	CTGI	T	TTAG	TTT					
+3	·V S		A	٧	Q	D	K	G	s	1	Н	K	F	<u> v</u>	<u>, P</u>	Y_	L,		<u> </u>	<u> </u>
1381	TGTC	TGGT	ĢÇ	TGTG	CAG	BAC	AAA	GGG'	TCAA	TC	CAC	LAAA	T	TGTC	CCT	TAC	CTG	ATT4	GCT	G
	3030	1001	CC	$\lambda \wedge \lambda \wedge$	<u>የርጥ</u> ረረ	TTG	ጥጥጥ	CCC	TTDA	AG	GTG'	$\mathbf{T}\mathbf{T}\mathbf{T}\mathbf{F}$	LΑ	ACAG	GGA	ATG	GAC	TAA	CGA	C
	ACAG	ACCA		MCMC																
+3	- G 1	Q	Н	3	С	Q	D	1	G	A	K	\$	Ļ	T	Q	<u> </u>	R	Α,	М	M -
+3 1441	GCAT	Q CCAA	H CA	S	C TGC	Q	D GAC	I ATT	G GGTG	A CC.	K AAG.	s AGCI	L T	T GACC	CAA	V GTC	R CGA	A GCC	M ATG	м - А
	- G 1	Q CCAA	H CA	S	C TGC	Q	D GAC	I ATT	G GGTG	A CC.	K AAG.	s AGCI	L T	T GACC	CAA	V GTC CAG	R CGA GCT	A GCC. CGG'	M ATG FAC	M· A T
	GCAT CGTA	Q CCAA GGTT 8	CA GT G	S CTCA GAGT E	C TGCC ACGC L	Q CAG GTC K	D GAC. CTG'	I ATT TAA	G GGTG CCAC K	CC. GG ¹ R	K AAG. TTC' T	s AGCT TCGA S	L TT AA S	T GACC CTGG A	Q CAA GTT Q	V GTC CAG V	R CGA GCT E	A GCC. CGG'	M ATG FAC G	M. A. T. V.
1441	GCAT CGTA M Y	Q CCAA GGTT S CTCT	CA GT G	S CTCA GAGT E	CATGCO	Q CAG GTC K	D GAC CTG	ATT TAA E GAG	GGTG CCAC K AAGA	GC. GG ^t R	K AAG. TTC' T ACG'	S AGCT FCGA S TCCT	L TT LA S	GACC CTGG A	CAA GTT Q CAG	V GTC CAG V GTG	R CGA GCT E GAA	GCC. CGG' G	M ATG FAC G GGC	M. A. T. V. G.
1441 +3	GCAT CGTA	Q CCAA GGTT S CTCT	CA GT G	S CTCA GAGT E	CATGCO	Q CAG GTC K	D GAC CTG	ATT TAA E GAG	GGTG CCAC K AAGA TTCT	GG'RGA	K AAG. TTC' T ACG'	S AGCT FCGA S TCCT AGGA	L TT LA S	GACC CTGG A	CAA GTT Q CAG	V GTC CAG V GTG	R CGA GCT E GAA	GCC. CGG' G	M ATG FAC G GGC	M. A. T. V. G.
1441 +3	GCAT CGTA M Y TGTA ACAT	Q CCAA GGTT S CTCT GAGA	GT GG GG CC L	GAGT GAGT E GGAG CCTC	C TGCC ACGC L CTTA CGAAT	Q CAG STC K AAG TTC Y	D GAC CTG' F TTT AAA	ATTO	GGTG CCAC K AAGA TTCT	A CC. GG. R GA. CT. L	K AAG. TTC' T ACG' TGC.	S AGCT PCGA S TCCT AGGA	L TT LA S TC LG	T GACC CTGG A AGCC TCGG	CAA GTT Q CAG	V GTC CAG V GTG CAC	GGA GCT E GAA CTT	A GCC. CGG' G GGT'	M TAC G G GGC CCG	M[A T V[G C
1441 +3 1501	GCAT CGTA M Y TGTA ACAT V H TCCA	Q CCAA GGTT S CTCT GAGA S	GG CC L	GAGT GAGT E GGAG CCTC H	CATGCC	Q CAG GTC K AAG PTC Y	D GAC CTG F TTT AAA E GAG	ATTO	GGTG CCAC K AAGA TTCT R	GC GG R GA CT L	K AAG TTC T ACG TGC	S AGCT FCGA S FCCT AGGA	L TT AA S CC AG	T GACC CTGG A AGCC TCGG	CAA GTT CAG GTC	V GTC CAG V GTG CAC	GGA GCT E GAA CTT	A GCC. CGG' GGT' CCA	M TAC G GGC CCG	M[ATV[GC A
1441 +3 1501 +3	GCAT CGTA M Y TGTA ACAT V H TCCA AGGT	CCAA GGTT 8 CTCT GAGA S TAGC	GT GG CC L	GAGT GAGT E GGAG CCTC H CCAT	C TGCC L CGAAT S TCGT	Q CAG STC K AAG TTC Y	GAC. CTG' F TTTC AAA. E GAG. CTC'	ATTO	GGTGCACKAAGATTCTACGGCGCGCGGCGGCGGCGGCGGCGGCGGGCG	GC GG F GA CT L	K AAG. TTC' ACG' TGC. F TTC'	S AGCT FCCT AGGA TGAT ACTA	L TT A S C C G	T GACC CTGG A AGCC TCGG	CAA GTT CAG GTC	V GTC CAG V GTG CAC	GGA GCT E GAA CTT TGA	A GCC. CGG' G GGTC CCA	M TAC G GGC CCG	M[ATV[GC AT
1441 +3 1501 +3	G I GCAT CGTA M Y TGTA ACAT V H TCCA AGGT	CCAA GGTT S CTCT GAGA S TAGC ATCG	GG GG CC L	S CTCA GAGT	CATGCO LACGAAT SGAAT SAGCA	Q CAG STC K AAG TTC Y FAT ATA	GAC. CTG' F TTTC AAA' E GAG. CTC' CCA	ATTO	GGTGCACKAAGATTCTACGGCGCGCCGCTAG	A CC GG R GA CT L TT AA	K AAG. TTC' TGC. F TTC' AAG.	S AGCT FCGA TCCT AGGA TGAT ACTA	L ST AA S CC AG AG AG AG	T GACC CTGG A AGCC TCGG TAGC ATCG AAAA	CAA GTT CAG GTC TCG AGC	V GTC CAG V GTG CAC ACA TGT	GGA GCT E GAA CTT TGA ACT	A GCC. GGT' CCA' TAA' ATT'	M TAC G GGC CCG GAT CTA	M[ATV[GC ATA
+3 1501 +3 1561	G I GCAT CGTA M Y TGTA ACAT V H TCCA AGGT CATT	Q CCAA GGTT 8 CTCT GAGA TAGC ATCG	GG GG CC L CT GA AG	GAGG GAGG E GGAG CCTC H CCAT GGTA	C TGCC L GAAT S AGC AGG C TGC	Q CAG STC K AAG TTC Y FAT AAA TTT	GAC. CTG' F TTTC AAA E GAG. CTC' CCAC	TAA EGAG CTC K AAG TTC	GGTGCACACACACACACACACACACACACACACACACACA	GG GG GA GT L TT AA AA TT.	K AAG. TTC' TGC. F TTC' AAG.	S AGCT PCGA S FCCT AGGA AGTG TCAC	L ST AS ST AG	T GACC CTGG A AGCC TCGG TAGC ATCG ATCG	CAA GTT CAG GTC TCG AGC	GTC CAG V GTG CAC ACA TGT GCT CGA	GAA GCTT GAA CTT TGA ACT TTA	A GCC. CGG' GGTCCA TAA TTTC	M TAC' G GGC CCG GAT CTA GTG	M[ATV[GC ATAT
+3 1501 +3 1561	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA	CCAA GGTT S CTCT GAGA S TAGC ATCG CATG	GG GG CT GA AG TC AT	GGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	C TGCC L GAAT S AGC AGG AC AGG AGG AGG AGG AGG AGG AGG	Q CAG GTC K AAG TTC Y ATA AAA TTT	GAC. CTG' F TTT' AAA' E GAG. CTC' CCA' GGT'	ATTO	G GGTG CCAC K AAGA TTCT R CGGC GCCG GCCG GTAG GATC	A CC GG F GA CT L TT AA AA TT AT	K AAG. TTC' TGC. AAG. TGC. AAG. TTGC	S AGCT S TCCT AGGA TGAT ACTA AGTC	L ST A S S A S A S A S A S A S A S A S A	T GACC CTGG A AGCC TCGG ATCG ATCG AAAA TTTT	CAA GTT CAG GTC TCG AGC AAT	V GTC CAG V GTG CAC ACA TGT GCT CGA	GAA CTT TGA ACT TTA AAT ATT	A GCC. CGG' GCA TAA ATT' AAA	MATG FAC GGC CCG GAT CTA GTG CAC	M.ATV[GC ATATC
1441 +3 1501 +3 1561 1621	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA AATT	CCAA GGTT S CTCT GAGA S TAGC ATCG CTAC TGTG	GG GG CC L CT GA AG TC TA	S CTCA GAGT E GGAG CCTC H CCAT GGTA TTTC AAAC GCTA CGAT	CATGCO LACGO S TTCGO AGCA CTGO ATTGO	Q CAG STC K AAG TTC Y ATA AAA TTT CTT GAA	D GAC. CTG' F TTTC AAA. E GAG. CTC' CCA. GGTC TAT' ATA.	TAAC GAG CTC K AAG TTC CAAC GTT TTG	GGTGCACKAAGATTCTAGGCCGGATCTGAAACTT	GG GG GA CT L TT AA AA TT AT	K AAG. TTC' TGC. AAG. TGC. ACG' TTG'	S AGCT PCGA S TCCT AGGA ACTA AGTG TCAC	L ST A S C A G A G A G A G A G A G A G A G A G A	T GACC CTGG A AGCC TCGG ATCG ATCG ATAAA ATTTT CTAT GATA	CAA GTT CAG GTC AGC AAT TTA	V GTC CAG V GTG CAC ACA TGT GCT CGA TTT	GAA CTT TGA ACT TTA AAT ATT	A GCC. CGG' GGT' CCA. TAA' ATT' AAA' TGT.	MATG. GGGCCGGATGCTACCACCACCACCACCACCACCACCACCACCACCACCAC	M.ATV[GC ATATCG
1441 +3 1501 +3 1561 1621	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA AATT	CCAA GGTT S CTCT GAGA S TAGC ATCG CTAC TGTG ACAC	GG GG CC L CT GA AG TC AT TA	S CTCA GAGT E GGAG CCTC H CCAT GGTA TTTC AAAC GCTA	CATGCO LACGAN SCTTI SCAN SCTCO LACGAN AGCI LACGAN L	Q CAG STC K AAG TTC Y FAT AAA TTT CTT GAA	D GAC CTG F TTTC AAA E GAG CTC CCA GGT TAT ATA	TAATTO KAAGTTC CAAAGTTTC CAAACTTAGAACTTAA	GGTGCACKAAGATTCTAGGCGGCGGATCTAGAAACTT	GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	K AAG. TTC' ACG' TTC' AAG. TGC. ACG' ACG'	S AGCT S TCCT AGGA TGAT ACTA TGAT ACTA TGAT	L ST A ST	T GACC CTGG A AGCC ATCG ATCG AAAA TTTTT CTAT GATA	CAA GTT CAG GTC TCG AGC AAT TTA	V GTC CAG V GTG CAC ACA TGT GCT CGA TTT AAA	GGA GCT GAA CTT TGA ACT TTA AAT ATT	A GCC. CGG' GGT' CCA' TAAATT' TTTT' AAA' TGT.	MATG. GGGCCGGGATGCTACCACACCACCACCACCACCACCACCACCACCACCACC	M[ATV[GC ATATCGT
1441 +3 1501 +3 1561 1621 1681	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA AATT TTAA ATTA	CCAA GGTT S CTCT GAGA TAGC ATCG CTAC TGTG ACAC TAAG	GG GG CC L CT GA AG TC GA	S CTCA GAGA GCTA CGTTA CTTCA CGTTA C	CATGCO CGAAT CGAAT AGCA CCTGT ATTGC	Q CAG FTC K AAG FTT ATA AAA FTT CTT AAA ACA FGT	D GAC CTG F TTTC AAA E GAG CTC CCA GGT TAT ATA AGT	I ATTO	GGTGCACKAAGATTCTAGGCGGCGATCAAACTTCAACGGTTGAACGTTTGAACAGAACGTTTGAACAGAACGTTTGAACAGAACAGAACAGAACAGAACAACAACAACAACAACA	GG	K AAG TTC' T ACG' F TTC' AAG TGC ACG' TTG' AACC CAA	S AGCT FCCT AGGA TGAT ACTA AGTG TGAT ACTA	L TT AA S TO AG	T GACC CTGG A AGCC ATCG AAAA TTTTT GATA AAGT	CAAA CTAG CAG CAG CTC CTC CTC CTC CTC CTC CTC CTC CTC CT	V GTC CAG V GTG CAC ACA TGT CGA TTT AAA TAT	GGA GCT E GAA CTT TGA ACT TTA AAT TAA GTT	A GCC. GGTCCA TAA ATT TTTCAAA TGT. ACA	MATG. GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	M[ATV[GC ATATCGTA
1441 +3 1501 +3 1561 1621 1681	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA AATT TTAA ATTA TAAT CAGG	CCAA GGTT S CTCT GAGA TAGC ATCG CTAC TGTG ACAC TAAG	H CA GG GG GG CC L CT GA AG CT CT GA GG	GGAGGGGTAGGGTAGGGTAGGGTAGGGTAGGGTAGGGT	CTGCCTCGCATCGCATCGCATCGCATCGCATCGCATCGC	Q CAG GTC K AAG TTC Y FATA AAA AAA CTT CTT GAA ACA FGT GGT	D GAC. CTG' F TTTCAAAA GAG. CTC' CCAC GGTC' TAT' ATA. AGT' TCA	ATTO GAG CTC K AAG TTC CAA GTT TTG AAC ATT	GGTGCACKAAGACTCAAACTCAACGCTGAACGCTAGCGCTAACGCTCAACGCAACCCAACCCAACCCAACCAA	GG	K AAG. TTC TACGTGC. AAG. TTCC AAG. TTGC AAC. GTTTGGAAC. GTT	S AGCT TCGA S TCCT AGGA ACTA AGTC AGGT ACTA ACTA ACTA	L TT LA S TO LG	T GACC CTGG A AGCC ATCG AAAA TTTT CTAT GATA AAGT TCTA	CAAC CAAC TTA TTA AAA CAA	V GTC CAG V GTG CAC ACA TGT GCT CGA TTT AAA TATA ATG	GGA GCT E GAA CTT TGA ACT TTA AAT TAA CTT CAA	A GCC. CGGG GGTCCCA TAAA TTTTT ACAA TCAAGT	MATG. ATG. GGGCCG GATCCTA CTA CAC AAC GTG GGTA ATG	M[ATV[GC ATATCGTAA
1441 +3 1501 +3 1561 1621 1681 1741	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA AATT TTAA ATTA TAAT CAGG GTCC	CCAA GGTT S CTCT GAGA S TAGC ATCG CTAC TGTG ACAC TAAG ATTC	H CA GG GG CC L CT GA AG TC GA GG GG CC	GGAGGGGAGGGGAGGGAGGGAGGGAGGGAGGGAGGGAG	C TACCO L COMMENT OF THE COMMENT OF	Q CAG K AAG TTC Y PATA AAAA TTT CTT GAA ACA TGT GGT CCA	GAC. CTG' F TTTT' AAA' E GAG. CTC' CCA' GGT' ATA AGT' ATA AGT' AAA	TAATTO	GGTGCACKAAGACTTGAAACTTGAACTTGAACTTGAACTTGAACTTGAACTTCG	GGA GAA AAA TT	K AAG. TTC' T ACG' TTC' AAG. TTC' AAG. TGC. ACG' TTG' AAC. CAA' GTT	S AGCT TCGA S TCCT AAGGA AAGTC AAGTC AAACC AAACC	L ST LA ST LG	T GACC CTGG A AGCC TCGG TAGC ATAGC ATAGT CTAT GATA AAGT TCTA AGAT	CAAA CAAA CAAA CGTT	V GTC CAG V GTG CAC ACA TGT CGA TTT AAA ATG TAC	GGA GCT GAA CTT TGA ACT TTA ATT TAA CTT TAA CTT	A GCC. CGGG GGTCCCA TAAA TTTTCAAAA TCAAAGT TCAAAGT TCAAAGT	MATGATAC GGATAC GATAC GATAC GATAC GATAC AAC GATAC GATAC AAC GATAC AAT CAAT AAT CAAT AAT CAAT AAT CAAT AAT	M[ATV[GC ATATCGTAAT
1441 +3 1501 +3 1561 1621 1681 1741	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA ATTA ATTA	CCAA GGTT S CTCT GAGA S TAGC ATCG CTAC TGTG ACAC TAAG ATTC GGGA CCCT	H GA GG GG L CT GA GG GG CT TT	GGAAACGGTATGTGAAAAC	CTTP	Q CAG K AAA AAA AAA AAA AAA AAA AAA AAA AAA	D GAC CTG' F TTTC AAA E GAG CTC' CCA GGT TAT ATA AGT TCA ATT AAA ATG	TAATTO TAA AATT TGA	GGTGCACKAAGATTCTAGGCAAACTTGAAACTTGAACCTTCGGCAA	GC GG R GA AA AA TT AA AA TT AA AA TT AA AA	K AAG TTC T ACG TGC TTC AAG TGC AAG TGC ACG TTG AAC CAA GTT GTA GGA GGC	S AGCT TCGA S TCGT AGGA AGTC AGTC AGTC AGTC AACT TTGAT ACT TGAT ACT TTGAT AC	L ST LA STOLE LA ST LA S	T GACC CTGG A AGCC ATCG AAAAA TTTTT CTAT GATA AAGT AGAT AAAAA AAAAA AAAAA AAAAA AAAAA AAAAA AAAA	CAAA CAAA CAAA CGTT	V GTC CAG V GTG CAC ACA ACA TGT TTT AAA ATG TAC AGG	GGA GCT GAA CTT TGA ACT TTAA ATT TAA CTT CAA TGG ACC	A GCC. CGGG GGTCCA TAAA TTTT AAAA TCA AGT TAGA TCA	MATG. ATG. GGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	M[ATV[GC ATATCGTAATA
1441 +3 1501 +3 1561 1621 1681 1741 1801	G I GCAT CGTA M Y TGTA ACAT TCCA AGGT CATT GTAA ATTA ATTA	CCAA GGTT S CTCT GAGA S TAGC ATCG CTAC TGTG ACAC TAAG ATTC GGGA CCCT AATG	H CA GT G GC L CT GA AG CC CT TA AG	GGAGGGGAGGGGAGGGGAGGGGAGGGGAGGGGAGGGGAGGGG	CATGCO LACGERAL SCHOOL CATGCO LACGERAL CATGCO	Q CAG K AAG PTC Y PATA AAA ACA ACA AAC PTG	D GAC. CTG' F TTTT' AAA' E GAG. CTC' CCA' GGT' ATA AGT' TCA ATT' AAA ATG' TAC	TAATTO TAA AATTO TAA AATTO AACTO TAA AATTO AACTO	GGTGCACKAAGACTTCAAGCTTCGAAGCTTCGGCAA	GCCGGGRGAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	K AAG TTC T ACG TGC TTC AAG TGC AAC GTT GTA GCAT GGC CCG	S AGCT TCGA TCGA TGAT AGCT TGAT ACT TGAT ACT TTGAT ACT T	L TT A S TO G G A T T G G G G	T GACC CTGG A AGCC TCGG TAGC ATCG AAAA TTTT CTAT GATA AAGT TCTA AGAT AAAG TTTC	CAAA GTT CGGAGGTC TCGGAGGAGGAGGAGGAGGAGGGGGGGGGG	V GTC CAG V GTG CAC ACA TGT GCT TAAA ATA ATA ATG AGG TCC	GGA GCT GAA CTT TGA ACT TTA ATT TAA GTT CAA TGG ACC	A GCC. GGCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	MATG. GATG. GATG. CTA CTA CTA CAC CAC CAC CAC CAC CAC CAC	M[ATV[GC ATATCGTAATAT
1441 +3 1501 +3 1561 1621 1681 1741 1801	G I GCAT CGTA ACAT TTAA ATTA TAAT CAGG GTCC TTTA	CCAA GGTT S CTCT GAGA TAGC ATCG CTAC TGTG ACAC TAAG ATTC GGGA CCCT AATG	H CA GT G GC L CT AA GG CC TT AA TT	GGAGGGGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	CATGCO ATGCO	Q CAG K AAG PTC Y ATA AAA PTT GAA ACA PGT GGT AAC PTT GAA CA AAC PTT GAA CA AAC PTT GAA CA AAC PTT GAA CA AAC PTT GAA AAC PTT GAA CA AAC AAC PTT GAA CA AAC PTT GAA CA AAC PTT GAA CA AAC PTT GAA CA AAC AAC PTT GAA CA AAC PT GAA CA AAC PT GAA CA AAC PT GAA CA AAC PT GAA CA AA	GAC. CTG' F TTTTAAA E GAG. CTC' CCA' GGT' ATA AGT' TCA ATG' TTCA	TAATTO CAAAGTTAAATTAAATTAAATTAAATTAAATTAAATTAA	GGTG CCAC K AAGA TTCT R CGGC GCTAG GATC TGAA ACTT CAAC GTTG AAGC TTCG GCAA CGTT	GCCGGGRGAAAAAAATTAAAATTTAAAATTTTAAAAATTTTTC	K AAG TTC' TGC. TTC' AAG TGC. ACG TGCAAC ACG ACG ACCCAAC GTT GGCACCCCCCCCCC	S AGCT TCGA S TCCTTGAT ACTA ACTA ACTA ACTA ACTA ACTA	L ST LA S CO L	T GACC CTGG A AGCC ATCG AAAA TTTT CAATA AAGT TCTA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAAGT TTTCA AAAAAGT TTTCA AAAAGT TTTCA AAAAGT TTTCA AAAAGT TTTCA AAAAGT TTTCA AAAAAGT TTTCA AAAAAGT TTTCA AAAAAGT TTTCA AAAAAGT TTTCA AAAAAAAAAA	CAAA CAAA CGTT TGC ACG TTAC CAAC CAAC CA	V GTC CAG V GTG CAC ACA TGT GCT TTT AAA ATG TAC AGG TCC CAT	GGA GCT GAA CTT TGA ACT TTA ATT TAA GTT CAA TGG ACC AAC	A GCC. GCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	MATG. GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	M[ATV[GC ATATCGTAATATC
1441 +3 1501 +3 1561 1621 1681 1741 1801 1861	G I GCAT CGTA ACAT TCCA AGGT TAAT TAAT CAGG GTCC TTTA AAAT AGGC TCCG	CCAA GGTT S CTCT GAGA TAGC CTAC CTAC CTAC CTAC TAAG ATTC GGGA ATTC CCCT CCCCT CCCCCCCCCC	H CA GT GG GC L CT AA CT GA GC TT AA CT AA	GGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	CTGCCTCCTCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	Q CAG K AAG TTC Y FATA AAA CTT GGT CCA AAC TTG AAA	GAC. CTG' F TTTCAAAACTCAAACTAACAACACAACACACACACACAC	TAATTO TAA AATTO TAAA AATTO TAAA AATTO CATA	GGTG CCAC K AAGA TTCT R CGGC GCTAG GATC TGAA ACTT GAAC GTTG AAGC TTCG GCAA CGTT AGGC TCCG	GCCGGGGACCTCAAAAAAAAAAAAAAAAAAAAAAAAAAA	K AAGA TTC T ACGGTGC AAG TTGC AACGTTGG AAC CAT GTA CAT GGC CGC GGC	SAGGITICGA STOCK SAGGITICGAT AGGITICGAT AGGITICGAT AGGITICAC AGGIT	L TT A S TO G C A T T C G C A T C C G C A T C C G C A T C C G C A T C C G C A T C C G C A T C C G C A T C C C C A T C C C C C A T C C C C	T GACC CTGG A AGCC ATCG AAAA TTTTT CTAT AGCT AAAGT TCTA AGAT AAAGT TCTA ACAT AAAGT AAAT A	CAA GTT CCAG GTC TCG AAT TTA CAA CCAA CC	V GTC CAG V GTG CAC ACA TGT TATA ATA ATG TAC AGG CAT GTA	GGA GCT GAA CTT TGA ACT TAA ATT CAA TGG ACC TTG CAC GTG	A GCC. GCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	MATG. GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	M[ATV[GC ATATCGTAATATCG
1441 +3 1501 +3 1561 1621 1681 1741 1801 1861	G I GCAT CGTA ACAT TTAA ATTA TAAT CAGG GTCC TTTA AAAT AGGC CACG	CCAA GGTT S CTCT GAGA S TAGC ATCG CTAC TGTG ACAC TGAG ATTC CCCT AATG CCCT CCCC CCCC	H CA GT GG CC L CT GA AC GG CC TT AA CT AA CT AA CT AA CT AA AC AC TT AA AC	GGAGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	C TGCC TAGCC TAGCC TAGCC TAGCC TGCC TGCC	CAGETC KAGETC Y PATA AAAA PETT GAAA PETT GEAA AACA PETT GEAA AACA PETT GEAA AACA PETT AAAA PETT AAAAA PETT AAAAA PETT AAAA PETT AAAAA PETT AAAAA PETT AAAAA PETT AAAAAAAAAA	GAC. CTG' F TTTT' AAA E GAG. CTC' CCA' GGT' ATA AGT' ATA ATG TTCA TTCA CGA	TAATTO K GAGGETTO CAA ATT TAA TAATT TGA ACT CATA ACT AACT A	GGTGCACKAAGACTGAAACTTGAACTTGAACTTGAACCTTCGCAACCCTCGAACCCTCGAACCCTCGAACCCTCGAACCCTCGAACCCTCGAACCCTCGAACCCCCAACCCCCAACCCCCAACCCCCAACCCCCAACCCC	GA GA AA TT AAA TT AAA TT TC AG CA	K AAG. TTC' F ACG. TTC' AAG. TGC. AAG. TGC. AAC. GTT. GTA GGC. GCG. GCG. GGGA	S AGCT TCGA S TCGT AGGA AGTC AGGA AGTC AAACC AAACC AAACC CCGGGC CCTAT	L TT A S C G G G G G G G G G G G G G G G G G G	T GACC CTGG A AGCC ATCG AAAA TTTT CAATA AAGT TCTA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAGT TTTCA AAAAGT TTTCA AAAAAGT TTTCA AAAAGT TTTCA AAAAGT TTTCA AAAAGT TTTCA AAAAGT TTTCA AAAAAGT TTTCA AAAAAGT TTTCA AAAAAGT TTTCA AAAAAGT TTTCA AAAAAAAAAA	CAAA GTT CGGGAAAT TAAA ACAA GTT GCGGGCCCAAAC CGTT CGGGCCCAAAC CGTT CGGGCCCAAC CGAGCCCAAC CTAC	V GTC CAG V GTG CAC ACA ACA TGT TATA ATA ATA ATA ATA CGT CGA CAT CGTA	GGA GCT GAA CTT TGA ACT TTA ATT TAA TGG ACC TTG CAC GTG	GCC. CGGG GGTCCCA TAAA TTTTAAAA TGTAAGT TAGGATCCGT AAAA TTTTTTTTTT	MATGATAC GGGCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	M[ATV[GC ATATCGTAATATCGC



Sheet 3 of 7

SELECTION SYSTEMS FOR GENETICALLY MODIFIED CELLS

HELLED CHOSES HARRIES & SECRETICES LLD



2041		CCTCGTGCGC GGAGCACGCG				
2101	CCTTTCTCCC	TTCGGGAAGC AAGCCCTTCG	GTGGCGCTTT	CTCAATGCTC	ACGCTGTAGG	TATCTCAGTT
2161	CGGTGTAGGT	CGTTCGCTCC GCAAGCGAGG	AAGCTGGGCT	GTGTGCACGA	ACCCCCCGTT	CAGCCCGACC
2221	GCTGCGCCTT	ATCCGGTAAC TAGGCCATTG	TATCGTCTTG	AGTCCAACCC	GGTAAGACAC	GACTTATCGC
2281		AGCCACTGGT TCGGTGACCA				
2341		GTGGTGGCCT CACCACCGGA				
2401	GAGACGACTT	GCCAGTTACC CGGTCAATGG	AAGCCTTTTT	CTCAACCATC	GAGAACTAGG	CCGTTTGTTT
2461	GGTGGCGACC	TAGCGGTGGT ATCGCCACCA	ААААААСААА	CGTTCGTCGT	CTAATGCGCG	TCTTTTTTTC
2521	CTAGAGTTCT	AGATCCTTTG TCTAGGAAAC	TAGAAAAGAT	GCCCCAGACT	GCGAGTCACC	TTGCTTTTGA
2581	GTGCAATTCC	GATTTTGGTC CTAAAACCAG	TACCGATCAA	TTAATTCGAC	GTTATTTGTT	AGTAATAAAA
2641	GTAACCTAGA	GTGTGTTGGT CACACAACCA	ÄAAAACACAC	CCGAACCCCC	TCCCCCTCCG	GTCTTACTGA
2701	GGTTCTCGAT	CAGGAAGGCA GTCCTTCCGT	CCAGTCTCTG	GGGTGACCTG	TTTGTCACCG	ACCTGAGACG
2761	TGGTATTGTG	ACAATCAACA TGTTAGTTGT	CCCCTCACTC	GACCTAGCTC	GATCTCAGGC	AATGTATTGA
2821	ATGCCATTTA	GGCCCGCCTG CCGGGCGGAC	CGACTGGCGG	GTTGCTGGGG	GCGGGTAACT	GCAGTTATTA
2881	CTGCATACAA	CCCATAGTAA GGGTATCATT	GCGGTTATCC	CTGAAAGGTA	ACTGCAGTTA	CCCACCTCAT
2941	AAATGCCATT	ACTGCCCACT TGACGGGTGA	ACCGTCATGT	AGTTCACATA	GTATACGGTT	CATGCGGGGG
3001	ATAACTGCAG	AATGACGGTA TTACTGCCAT	TTACCGGGCG	GACCGTAATA	CGGGTCATGT	ACTGGAATAC
3061	CCTGAAAGGA	ACTTGGCAGT TGAACCGTCA	TGTAGATGCA	TAATCAGTAG	CGATAATGGT	ACCACTACGC
3121	CAAAACCGTC	TACATCAATG ATGTAGTTAC	CCGCACCTAT	CGCCAAACTG	AGTGCCCCTA	AAGGTTCAGA
3181	GGTGGGGTAA	GACGTCAATG CTGCAGTTAC	CCTCAAACAA	AACCGTGGTT	TTAGTTGCCC	TGAAAGGTTT
3241	TACAGCATTG	AACTCCGCCC TTGAGGCGGG	GTAACTGCGT	TTACCCGCCA	TCCGCACATG	CCACCCTCCA
3301	GATATATTCG	AGAGCTCGTT TCTCGAGCAA	ATCACTTGGC	AGTCTAGCGG	ACCTCTGCGG	TAGGTGCGAC
3361	TTTTGACCTC AAAACTGGAG	CATAGAAGAC GTATCTTCTG	ACCGGGACCG TGGCCCTGGC	ATCCAGCCTC TAGGTCGGAG	GCGCCGGCCC	TTGCCACGTA



HELLER EHRMAN WHITE & MCAULIFFE LLP Sheet 4 of 7

SELECTION SYSTEMS FOR GENETICALLY MODIFIED CELLS

DOCKET NO. 24751-2502 Applicant: Jensen Filed: April 30, 2001

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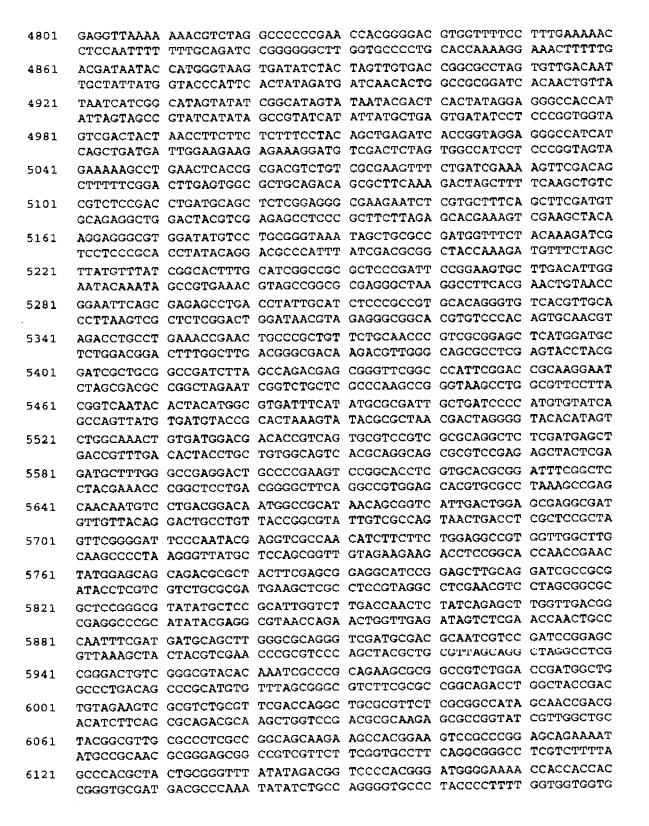


3421		ATTCCCCGTG TAAGGGGCAC				
3481	CACCCCCTTG	GCTTCTTATG CGAAGAATAC	CATGCTATAC	TGTTTTTGGC	TTGGGGTCTA	TACACCCCCG
3541	CTTCCTCATG	TTATAGGTGA AATATCCACT	TGGTATAGCT	TAGCCTATAG	GTGTGGGTTA	TTGACCATTA
3601	TTGACCACTC	CCCTATTGGT	GACGATACTT	TCCATTACTA	ATCCATAACA	TGGCTCTTTG
3661	CCACAACTCT	GGGATAACCA CTTTATTGGC	TATATGCCAA	TACACTGTCC	TTCAGAGACT	GACACGGACT
3721	CTGTATTTTT	GAAATAACCG ACAGGATGGG	GTCTCATTTA	TTATTTACAA	ATTCACATAT	ACAACACCAC
3781	CGTCCCCAGT	TGTCCTACCC GCCCGCAGTT	TTTATTAAAC	ATAACGTGGG	ATCTCCACGC	GAATCTCGGG
3841	TACGTGTTCC	CGGGCGTCAA GGACATGGGC	TCTTCTCCGG	TAGCGGCGGA	GCTTCTACAT	CCGAGCCCTG
3901		CCTGTACCCG TCCAGCGACT				
3961	GAGGGTACGG	AGGTCGCTGA CACAGCACGA	GTACCAGCGA	GCCGTCGAGG	AACGAGGATT	GTCACCTCCG
	GTCTGAATCC	GTGTCGTGCT TCTGAAAATG	ACGGGTGGTG	GTGGTCACAC	GGCGTGTTCC	GGCACCGCCA
4021	TCCCATACAC	AGACTTTTAC GCGGCAGAAG	TCGAGCCCCT	CGCCCGAACG	TGGCGACTGC	GTAAACCTTC
4081	TGAATTCCGT	CGCCGTCTTC	TTCTACGTCC	GTCGACTCAA	CAACACAAGA	CTATTCTCAG
4141	AGAGGTAACT	CCCGTTGCGG	TGCTGTTAAC ACGACAATTG	CCACCTCCCG	TCACATCAGA	CTCGTCATGA
4201	CGTTGCTGCC	GCGCGCGCCA CGCGCGCGGT	CCAGACATAA	TAGCTGACAG ATCGACTGTC	ACTAACAGAC TGATTGTCTG	TGTTCCTTTC ACAAGGAAAG
	CGTTGCTGCC GCAACGACGG	GCGCGCGCCA CGCGCGCGGT	CCAGACATAA GGTCTGTATT	TAGCTGACAG ATCGACTGTC MCS	TGATTGTCTG	ACAAGGAAAG
4201 4261	CGTTGCTGCC GCAACGACGG CATGGGTCTT	GCGCGCGCCA CGCGCGCGGT TTCTGCAGTC	CCAGACATAA GGTCTGTATT ACCCGGGGGA	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC	TGATTGTCTG GTAGCTCTAG	ACAAGGAAAG ATTGAGTCGA
	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC	GCGCGCGCA CGCGCGCGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG	TGATTGTCTG GTAGCTCTAG CATCGAGATC TTTGTCTATA	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC
4261	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG	GCGCGCGCA CGCGCGCGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT GCTTCGGCGA	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG ACCTTATTCC	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCACACGC	TGATTGTCTG GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG
4261	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG CACCATATTG	GCGCGCGCA CGCGCGCGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT GCTTCGGCGA CCGTCTTTTG	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG ACCTTATTCC GCAATGTGAG	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCACACGC GGCCCGGAAA	TGATTGTCTG GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT CCTGGCCCTG	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG TCTTCTTGAC
4261 4321	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG CACCATATTG GTGGTATAAC GAGCATTCCT	CGCGCGCCA CGCGCGCGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT GCTTCGGCGA CCGTCTTTTG GGCAGAAAAC AGGGGTCTTT	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG ACCTTATTCC GCAATGTGAG CGTTACACTC	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCACACGC GGCCCGAAA CCGGGCCTTT CAAAGGAATG	TGATTGTCTG GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT CCTGGCCCTG GGACCGGGAC CAAGGTCTGT	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG TCTTCTTGAC AGAAGAACTG TGAATGTCGT
4261 4321 4381	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG CACCATATTG GTGGTATAAC GAGCATCCT CTCGTAAGGA	GCGCGCGCA CGCGCGCGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT GCTTCGGCGA CCGTCTTTTG GGCAGAAAAC AGGGGTCTTT TCCCCAGAAA	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG ACCTTATTCC GCAATGTGAG CGTTACACTC CCCCTCTCGC GGGGAGAGCG AAGCTTCTTG	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCACACGC GGCCCGGAAA CCGGGCCTTT CAAAGGAATG GTTTCCTTAC AAGACAACA	TGATTGTCTG GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT CCTGGCCTG GGACCGGGAC CAAGGTCTGT GTTCCAGACA ACGTCTGTAG	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG TCTTCTTGAC AGAAGAACTG TGAATGTCGT ACTTACAGCA CGACCCTTTG
4261 4321 4381 4441	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG CACCATATTG GTGGTATAAC GAGCATTCCT CTCGTAAGGA GAAGGAAGCA CTTCCTTCGT	GCGCGCGCA CGCGCGCGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT GCTTCGGCGA CCGTCTTTTG GGCAGAAAAC AGGGGTCTTT TCCCCAGAAA GTTCCTCTGG CAAGGAGACC	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG ACCTTATTCC GCAATGTGAG CGTTACACTC CCCCTCTCGC GGGGAGAGCG AAGCTTCTTG TTCGAAGAAC CTGGCGACAG	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCCGGAAA CCGGGCCTTT CAAAGGAATG GTTTCCTTAC AAGACAACA TTCTGTTTGT GTGCCTCTGC	GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT CCTGGCCCTG GGACCGGGAC CAAGGTCTGT GTTCCAGACA ACGTCTGTAG TGCAGACATC GGCCAAAAGC	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG TCTTCTTGAC AGAAGAACTG TGAATGTCGT ACTTACAGCA CGACCCTTTG GCTGGGAAAC CACGTGTATA
4261 4321 4381 4441 4501	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG CACCATATTG GTGGTATAAC GAGCATTCCT CTCGTAAGGA GAAGGAAGCA CTTCCTTCGT CAGGCAGCGG GTCCGTCGCC	CGGGGGGCA CGCGCGGGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT GCTTCGGCGA CCGTCTTTTG GGCAGAAAAC AGGGGTCTTT TCCCCAGAAA GTTCCTCTGG CAAGGAGACC AACCCCCAC TTGGGGGGTG GCAAAGGCGG	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCT TGGAATAAGG ACCTTATTCC GCAATGTGAG CGTTACACTC CCCCTCTCGC GGGGAGAGCG AAGCTTCTTG TTCGAAGAAC CTGGCGACAG GACCGCTGTC CACAACCCCA	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCACACGC GGCCCGGAAA CCGGGCCTTT CAAAGGAATG GTTTCCTTAC AAGACAAACA TTCTGTTTGT GTGCCTCTGC CACGGAGACG GTGCCACGTT	GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT CCTGGCCCTG GGACCGGGAC CAAGGTCTGT GTTCCAGACA ACGTCTGTAG TGCAGACATC GGCCAAAAGC CCGGTTTTCG GTGAGTTTCG	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG TCTTCTTGAC AGAAGAACTG TGAATGTCGT ACTTACAGCA CGACCCTTTG GCTGGGAAAC CACGTGTATA GTGCACATAT TAGTTGTGA
4261 4321 4381 4441 4501 4561	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG CACCATATTG GTGGTATAAC GAGCATTCCT CTCGTAAGGA CTTCCTTCGT CAGGCAGCGG GTCCGTCGCC AGATACACT TCTATGTGGA	GCGCGCGCA CGCGCGCGT TTCTGCAGTC AAGACGTCAG CGAAGCCGCT GCTTCGGCGA CCGTCTTTTG GGCAGAAAAC AGGGGTCTTT TCCCCAGAAA GTTCCTCTGG CAAGGAGACC AACCCCCAC TTGGGGGGTG GCAAAGGCGG CGTTTCCGCC	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG ACCTTATTCC GCAATGTGAG CGTTACACTC CCCCTCTCGC GGGGAGAGCG AAGCTTCTTG TTCGAAGAAC CTGGCGACAG GACCGCTGTC CACAACCCCA GTGTTGGGGT CAAGCGTATT	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCACACGC GGCCCGGAAA CCGGGCCTTT CAAAGGAATG GTTTCCTTAC AAGACAAACA TTCTGTTTGT GTGCCTCTGC CACGGAGACG GTGCCACGTT CACGGTGCAA CAACAAGGGG	GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT CCTGGCCCTG GGACCGGGAC CAAGGTCTGT GTTCCAGACA ACGTCTGTAG TGCAGACATC GGCCAAAAGC CCGGTTTTCG GTGAGTTGGA CACTCAACCT CTGAAGGATG	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG TCTTCTTGAC AGAAGAACTG TGAATGTCGT ACTTACAGCA CGACCCTTTG GCTGGGAAAC CACGTGTATA GTGCACATAT TAGTTGTGGA ATCAACACCT CCCAGAAGGT
4261 4321 4381 4441 4501 4561 4621	CGTTGCTGCC GCAACGACGG CATGGGTCTT GTACCCAGAA CGTTACTGGC GCAATGACCG CACCATATTG GTGGTATAAC GAGCATTCCT CTCGTAAGGA CTTCCTTCGT CAGGCAGCGC GTCCGTCGCC AGATACACCT TCTATGTGGA AAGAGTCAAA TTCTCAGTTT	CCGCGCGCCA CGCGCGCGT TTCTGCAGTC AAGACCGCT GCTTCGGCGA CCGTCTTTTG GGCAGAAAAC AGGGGTCTTT TCCCCAGAAA GTTCCTCTGG CAAGGAGACC AACCCCCAC TTGGGGGGTG GCAAAGGCGG CGTTTCCGCC TGGCTCTCCT ACCGAGAGGA ATGGGATCTG	CCAGACATAA GGTCTGTATT ACCCGGGGGA TGGGCCCCCT TGGAATAAGG ACCTTATTCC GCAATGTGAG CGTTACACTC CCCCTCTCGC GGGGAGAGCG AAGCTTCTTG TTCGAAGAAC CTGGCGACAG GACCGCTGTC CACAACCCCA GTGTTGGGGT CAAGCGTATT GTTCGCATAA ATCTGGGGCC	TAGCTGACAG ATCGACTGTC MCS TCCTTCGAAC AGGAAGCTTG CCGGTGTGCG GGCCCGGAAA CCGGGCCTTT CAAAGGAATG GTTTCCTTAC AAGACAACA TTCTGTTTGT GTGCCTCTGC CACGGAGACG GTGCCACGTT CACGGTGCAA CAACAAGGGG GTTGTTCCCC TCGGTGCACA	GTAGCTCTAG CATCGAGATC TTTGTCTATA AAACAGATAT CCTGGCCCTG GGACCGGGAC CAAGGTCTGT GTTCCAGACA ACGTCTGTAG TGCAGACATC GGCCAAAAGC CCGGTTTTCG GTGAGTTGGA CACTCAACCT CTGAAGGATG GACTTCCTAC TGCTTTACAT	ACAAGGAAAG ATTGAGTCGA TAACTCAGCT TGTTATTTTC ACAATAAAAG TCTTCTTGAC AGAAGAACTG TGAATGTCGT ACTTACAGCA CGACCCTTTG GCTGGGAAAC CACGTGTATA GTGCACATAT TAGTTGTGA ATCAACACCT CCCAGAAGGT GGGTCTTCCA



SELECTION SYSTEMS FOR GENETICALLY MODIFIED CELLS







Sheet 6 of 7 SELECTION SYSTEMS FOR GENETICALLY MODIFIED CELLS DOCKET NO. 24354 2502

DOCKET NO. 24751-2502

Applicant: Jensen Filed: April 30, 2001

6181	GCAACTGCTG	GTGGCCCTGG CACCGGGACC	GTTCGCGCGA CAAGCGCGCT	CGATATCGTC GCTATAGCAG	TACGTACCCG ATGCATGGGC	AGCCGATGAC TCGGCTACTG
co.4.1		GTGCTGGGGG				
6241	AATGACCGCC	CACGACCCCC	GAAGGCTCTG	TTAGCGCTTG	TAGATGTGGT	GTGTTGTGGC
6301		GGTGAGATAT				
	GGAGCTGGTC	CCACTCTATA	GCCGGCCCCT	GCGCCGCCAC	CATTACTGTT	CGCGGGTCTA
6361	AACAATGGGC	ATGCCTTATG	CCGTGACCGA	CGCCGTTCTG	GCTCCTCATA	TCGGGGGGGA
	TTGTTACCCG	TACGGAATAC	GGCACTGGCT	GCGGCAAGAC	CGAGGAGTAT	AGCCCCCCT
6421		TCACATGCCC				
0421	CCCACCCTCC	AGTGTACGGG	eceeeeecee	GGAGTGGGAG	TAGAAGCTGG	CGGTAGGGTA
		CTGTGCTACC				
6481	CGCCGCCCTC	GACACGATGG	CGGCCGCGCG	CAMOCAAMAC	CCCTCCTACT	CCCCCCTCCC
6541	CGTGCTGGCG	TTCGTGGCCC	TCATCCCGCC	GACCTTGCCC	GGCACCAACA	ACCACCA ACC
		AAGCACCGGG				
6601	GGCCCTTCCG	GAGGACAGAC	ACATCGACCG	CCTGGCCAAA	CGCCAGCGCC	CCGCCGAGCG
		CTCCTGTCTG				
6661	GCTGGACCTG	GCTATGCTGG	CTGCGATTCG	CCGCGTTTAC	GGGCTACTTG	CCAATACGGT
	CGACCTGGAC	CGATACGACC	GACGCTAAGC	GGCGCAAATG	CCCGATGAAC	GGTTATGCCA
6721		CAGTGCGGCG				
• • • •	CGCCATAGAC	GTCACGCCGC	CCAGCACCGC	CCTCCTGACC	CCTGTCGAAA	GCCCCTGCCG
6781		CAGGGTGCCG				
0/61	CCACCCCCC	GTCCCACGGC	TCGGGGTCTC	GTTGCGCCCG	GGTGCTGGGG	TATAGCCCCT
		ACCCTGTTTC				
6841	CACGTTATTT	TGGGACAAAG	CCCCCCCCC	CAACGACCGG	GGGTTGCCGC	TGGACATATT
		TGGGCCTTGG				
6901	CGTGTTTGCC	ACCCGGAACC	MCGICIIGGC	CHARCGCCIC	CCAAGGTACG	TGCAGAAATA
6961	CCTGGATTAC	GACCAATCGC	CCGCCGGCTG	CCGGGACGCC	CIGCIGCAAC	AATGGAGGCC
		CTGGTTAGCG				
7021	GATGGTCCAG	ACCCACGTCA	CCACCCCGG	CTCCATACCG	ACGATATGCG	MCCTGGCGCG
		TGGGTGCAGT				
7081	CACGTTTGCC	CGGGAGATGG	GGGAGGCTAA	CTGAGTCGAG	AATTCGCTAG	AGGGCCCTAT
	GTGCAAACGG	GCCCTCTACC	CCCTCCGATT	GACTCAGCTC	TTAAGCGATC	TCCCGGGATA
7141	TCTATAGTGT	CACCTAAATG	CTAGAGCTCG	CTGATCAGCC	TCGACTGTGC	CTTCTAGTTG
	AGATATCACA	GTGGATTTAC	GATCTCGAGC	GACTAGTCGG	AGCTGACACG	GAAGATCAAC
7201	CCAGCCATCT	GTTGTTTGCC	CCTCCCCCGT	GCCTTCCTTG	ACCCTGGAAG	GTGCCACTCC
. •	GGTCGGTAGA	CAACAAACGG	GGAGGGGGCA	CGGAAGGAAC	TGGGACCTTC	CACGGTGAGG
7261	CACTGTCCTT	ТССТААТААА	ATGAGGAAAT	TGCATCGCAT	TGTCTGAGTA	GGTGTCATTC
7201	GTGACAGGAA	AGGATTATTT	TACTCCTTTA	ACGTAGCGTA	ACAGACTCAT	CCACAGTAAG
7321	TATTCTCCCC	CATCCCCTCC	GGCAGGACAG	CAAGGGGGAG	GATTGGGAAG	ACAATAGCAG
7,321	DTAAGACCCC	CCACCCCACC	CCGTCCTGTC	GTTCCCCCTC	CTAACCCTTC	TGTTATCGTC
7201	ATAMONOCOLO	GGCCCAATTG	CTCGAGCGGC	CGCAATAAAA	TATCTTTATT	TTCATTACAT
7381	CCMACCCCAG	CCGGGTTAAC	GAGCTCGCCG	GCGTTATTTT	ATAGAAATAA	AAGTAATGTA
	CGIACGCGIC	GTTTTTTGTG	መር እ አመርር መኦ እ	СТАВСВТВСС	CTCTCCATCA	AAACAAAACG
7441	CTGTGTGTTG	CAAAAAACAC	TOWATCO LAW	CATTCTATAC	GAGAGGTAGT	TTTGTTTTGC
	GACACACAAC	CAAAAAACAC	MULINGONII	COCA CHOCA A	CHCCACCTCC	CAGAACATTT
7501	AAACAAAACA	AACTAGCAAA	ATAGGCTGTC	CCCAGIGCAA	CACCACCACC	GTCTTGTAAA
	TTTGTTTTGT	TTGATCGTTT	TATCCGACAG	GOGICACGII	CVCGICCVCG	



HELLEH EHRMAN WHITE & MCAULIFFE LLP Sheet 7 of 7 SELECTION SYSTEMS FOR GENETICALLY MODIFIED CELLS



7561		GGATCTGCGA				
	GAGATAGCTT	CCTAGACGCT	AGCGAGGCCA	CGGGCAGTCA	CCCGTCTCGC	GTGTAGCGGG
7621	ACAGTCCCCG	AGAAGTTGGG	GGGAGGGGTC	GGCAATTGAA	CCGGTGCCTA	GAGAAGGTGG
	TGTCAGGGGC	TCTTCAACCC	CCCTCCCCAG	CCGTTAACTT	GGCCACGGAT	CTCTTCCACC
7681	CGCGGGGTAA	ACTGGGAAAG	TGATGTCGTG	TACTGGCTCC	GCCTTTTTCC	CGAGGGTGGG
	GCGCCCCATT	TGACCCTTTC	ACTACAGCAC	ATGACCGAGG	CGGAAAAAGG	GCTCCCACCC
7741	GGAGAACCGT		AGTAGTCGCC			
	CCTCTTGGCA	TATATTCACG	TCATCAGCGG	CACTTGCAAG	AAAAAGCGTT	GCCCAAACGG
7801	GCCAGAACAC	AGCTGAAGCT	TCGAGGGGCT	CGCATCTCTC	CTTCACGCGC	CCGCCGCCCT
	CGGTCTTGTG	TCGACTTCGA	AGCTCCCCGA	GCGTAGAGAG	GAAGTGCGCG	GGCGGCGGGA
7861	ACCTGAGGCC		CCGGTTGAGT			
	TGGACTCCGG	CGGTAGGTGC	GGCCAACTCA	GCGCAAGACG	GCGGAGGGCG	GACACCACGG
7921		CGTCCGCCGT				
	AGGACTTGAC	GCAGGCGGCA	GATCCATTCA	AATTTCGAGT	CCAGCTCTGG	CCCGGAAACA
7981		CTTGGAGCCT				
		GAACCTCGGA				
8041		TCTACGTCTT				
	GAACGAGTTG	AGATGCAGAA	ACAAAGCAAA	AGACAAGACG	CGGCAATGTC	TAGGTTCGAC
8101	TGACCGGCGC	CTACGTAAGT			AAAAGAGTGT	
	ACTGGCCGCG	GATGCATTCA	CTATAGATGA	TCTAAATAGT	TTTTCTCACA	ACTGAACACT
8161	GCGCTCACAA	TTGATACTTA			TCGACTACTA	
	CGCGAGTGTT	AACTATGAAT	CTAAGTAGCT	CTCCCTGTGC	AGCTGATGAT	TGGAAGAAGA
8221	CTTTCCTACA	GCTGAGAT				
	GAAAGGATGT	CGACTCTA				